Serial Number: 10/084,553

CRF Processing Date: 7/31/02
Edited by: _____ DC
Verified by: _____ DC _____ (STIC staff)

ENTERED

- [ ] Changed a file from non-ASCII to ASCII

- [ ] Changed the margins in cases where the sequence text was "wrapped" down to the next line.

- [ ] Edited a format error in the Current Application Data section, specifically: _____

- [ ] Edited the Current Application Data section with the actual current number. The number inputted by the applicant was [ ] the prior application data; or [ ] other _____.

- [ ] Added the mandatory heading and subheadings for "Current Application Data".

- [ ] Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.

- [ ] Changed the spelling of a mandatory field (the headings or subheadings), specifically: _____

- [ ] Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: _____

- [ ] Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: _____

- [ ] Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.

- [ ] Inserted colons after headings/subheadings. Headings edited included: _____

- [ ] Deleted extra, invalid, headings used by an applicant, specifically: _____

- [ ] Deleted: [ ] non-ASCII "garbage" at the beginning/end of files; [ ] secretary initials/filename at end of file; [ ] page numbers throughout text; [ ] other invalid text, such as_____.

- [✓] Inserted mandatory headings, specifically: L2207 in Seq. 2

- [ ] Corrected an obvious error in the response, specifically: _____

- [ ] Edited identifiers where upper case is used but lower case is required, or vice versa.

- [ ] Corrected an error in the Number of Sequences field, specifically: _____

- [ ] A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.

- [ ] Deleted *ending* stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____

- [ ] Other: _____

*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95

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           RAW SEQUENCE LISTING                   DATE: 07/31/2002
           PATENT APPLICATION:  US/10/084,553     TIME: 12:41:33

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      4         Wang, Zhi-Yong
      5         Sun, Lin
      6         The Regents of the University of California
      8 <120> TITLE OF INVENTION: Phytochrome Regulated Transcription Factor for Control
      9         of Higher Plant Development
     11 <130> FILE REFERENCE: 023070-124200US
     13 <140> CURRENT APPLICATION NUMBER: US 10/084,553
C--> 14 <141> CURRENT FILING DATE: 2002-07-09
     16 <150> PRIOR APPLICATION NUMBER: US 08/843,572
     17 <151> PRIOR FILING DATE: 1997-04-18
     19 <160> NUMBER OF SEQ ID NOS: 27
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 91 tattagggag cagctgaatc aaaggaggaa gaagaagaag aagagccttt ttgaggccat 240
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129 ggagcttagt g atg gag aca aat tcg tct gga gaa gat ctg gtt att aag 1370
130             Met Glu Thr Asn Ser Ser Gly Glu Asp Leu Val Ile Lys
131              1               5                   10
133 gtaaattaac taaattttag ggggaagatg attgttttag gtgtcaaaga ttgagaattt 1430
135 taatgaaact tgatatag act cgg aag cca tat acg ata aca aag caa cgt 1481
136             Thr Arg Lys Pro Tyr Thr Ile Thr Lys Gln Arg
137                     15                  20
139 gaa agg tgg act gag gaa gaa cat aat aga ttc att gaa gct ttg agg 1529
140 Glu Arg Trp Thr Glu Glu Glu His Asn Arg Phe Ile Glu Ala Leu Arg
141 25              30                  35                  40
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143 ctt tat ggt aga gca tgg cag aag att gaa g gttgattttt atttcccttt   1580
144 Leu Tyr Gly Arg Ala Trp Gln Lys Ile Glu
145                 45                      50
147 atatgtctta ttttttgtgt ttgcagaggt ttgtcttcaa actgatttgc tttttttcat  1640
149 ttggacag  aa cat gta gca aca aaa act gct gtc cag ata aga agt cac  1689
150         Glu His Val Ala Thr Lys Thr Ala Val Gln Ile Arg Ser His
151                         55                  60
153 gct cag aaa ttt ttc tcc aag gtaaaatcgg ttaattttga aatgatgttc      1740
154 Ala Gln Lys Phe Phe Ser Lys
155 65                70
157 tcatcttcat tggcttaatg cttaagactt attgaaagcc aggcaagttt tctgcttctt  1800
159 ttgcttctta gtcaggagat agatagatta cgtttttaga gtttagtaat gagcaataag  1860
161 tcttaaaata gttggagaaa tgacgagatg taatcgtttt cttttgttta tgcctatatc  1920
163 ttgttaatcc acaaacatgt acatagattc ttcagaagaa tgttagtttc tttagattct  1980
165 tcagataaac ttgtgtcttc ttaccgattc tgaggtagtg gcaaagtgg gctgagtgct   2040
167 agaaattttt gaatgttcct tgtgataagc catagaggta aaccattttt gattttccag  2100
169 ttctgtcatt taaacttgtt aggtgtcatt agatttttgt ttgtttacgt ttgtttagag  2160
171 ggtaacaaaa ctactctcat ctctctcag gta gag aaa gag gct gaa gct aaa   2213
172                              Val Glu Lys Glu Ala Glu Ala Lys
173                                              75
175 ggt gta gct atg ggt caa gcg cta gac ata gct att cct cct cca cgg   2261
176 Gly Val Ala Met Gly Gln Ala Leu Asp Ile Ala Ile Pro Pro Pro Arg
177 80                  85                  90                  95
179 cct aag cgt aaa cca aac aat cct tat cct cga aag acg gga agt gga   2309
180 Pro Lys Arg Lys Pro Asn Asn Pro Tyr Pro Arg Lys Thr Gly Ser Gly
181                 100                 105                 110
183 acg atc ctt atg tca aaa acg ggt gtg aat gat gga aaa gag tcc ctt   2357
184 Thr Ile Leu Met Ser Lys Thr Gly Val Asn Asp Gly Lys Glu Ser Leu
185                 115                 120                 125
187 gga tca gaa aaa gtg tcg cat cct gag gtgattttca tggtcatatg         2404
188 Gly Ser Glu Lys Val Ser His Pro Glu
189         130                 135
191 gcatcttttt gcagtgtgtc acattgctcc tcatgttatt aatacagatt gtgtgcttcg  2464
193 tttatag atg gcc aat gaa gat cga caa caa tca aag cct gaa gag aaa   2513
194         Met Ala Asn Glu Asp Arg Gln Gln Ser Lys Pro Glu Glu Lys
195                 140                 145                 150
197 act ctg cag gaa gac aac tgt tca gat tgt ttc act cat cag tat ctc   2561
198 Thr Leu Gln Glu Asp Asn Cys Ser Asp Cys Phe Thr His Gln Tyr Leu
199                 155                 160                 165
201 tct gct gca tcc tcc atg aat aaa agt tgt ata gag aca tca aac gca   2609
202 Ser Ala Ala Ser Ser Met Asn Lys Ser Cys Ile Glu Thr Ser Asn Ala
203                 170                 175                 180
205 agc act ttc cgc gag ttc ttg cct tca cgg gaa gag gtaaaaaaca        2655
206 Ser Thr Phe Arg Glu Phe Leu Pro Ser Arg Glu Glu
207         185                 190
209 atctttcatt gctatttgag gttttaagac gattagtact tttcatgaaa ctaaaaccgt  2715
211 gggggaataa cag gga agt cag aat aac agg gta aga aag gag tca aac     2764
212             Gly Ser Gln Asn Asn Arg Val Arg Lys Glu Ser Asn
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219 cag act tat ccg atg cat atc cct gtg cta gtg cca ttg ggg agc tca     2860
220 Gln Thr Tyr Pro Met His Ile Pro Val Leu Val Pro Leu Gly Ser Ser
221         225             230             235
223 ata aca agt tct cta tca cat cct cct tca gag cca gat agt cat ccc     2908
224 Ile Thr Ser Ser Leu Ser His Pro Pro Ser Glu Pro Asp Ser His Pro
225     240             245             250
227 cac aca gtt gca gga gat tat cag tcg ttt cct aat cat ata atg tca     2956
228 His Thr Val Ala Gly Asp Tyr Gln Ser Phe Pro Asn His Ile Met Ser
229 255             260             265             270
231 acc ctt tta caa aca ccg gct ctt tat act gcc gca act ttc gcc tca     3004
232 Thr Leu Leu Gln Thr Pro Ala Leu Tyr Thr Ala Ala Thr Phe Ala Ser
233             275             280             285
235 tca ttt tgg cct ccc gat tct agt ggt ggc tca cct gtt cca ggg aac     3052
236 Ser Phe Trp Pro Pro Asp Ser Ser Gly Gly Ser Pro Val Pro Gly Asn
237             290             295             300
239 tca cct ccg aat ctg gct gcc atg gcc gca gcc act gtt gca gct gct     3100
240 Ser Pro Pro Asn Leu Ala Ala Met Ala Ala Ala Thr Val Ala Ala Ala
241         305             310             315
243 agt gct tgg tgg gct gcc aat gga tta tta cct tta tgt gct cct ctt     3148
244 Ser Ala Trp Trp Ala Ala Asn Gly Leu Leu Pro Leu Cys Ala Pro Leu
245     320             325             330
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248 Ser Ser Gly Gly Phe Thr Ser His Pro Pro Ser Thr Phe Gly Pro Ser
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251 tgt gat gta gag tac aca aaa gca agc act tta caa cat ggt tct gtg     3244
252 Cys Asp Val Glu Tyr Thr Lys Ala Ser Thr Leu Gln His Gly Ser Val
253             355             360             365
255 cag agc cga gag caa gaa cac tcc gag gca tca aag gct cga tct tca     3292
256 Gln Ser Arg Glu Gln Glu His Ser Glu Ala Ser Lys Ala Arg Ser Ser
257             370             375             380
259 ctg gac tca gag gat gtt gaa aat aag agt aaa cca gtt tgt cat gag     3340
260 Leu Asp Ser Glu Asp Val Glu Asn Lys Ser Lys Pro Val Cys His Glu
261         385             390             395
263 cag cct tct gca aca cct gag agt gat gca aag ggt tca gat gga gca     3388
264 Gln Pro Ser Ala Thr Pro Glu Ser Asp Ala Lys Gly Ser Asp Gly Ala
265     400             405             410
267 gga gac aga aaa caa gtt gac cgg tcc tcg tgt ggc tca aac act ccg     3436
268 Gly Asp Arg Lys Gln Val Asp Arg Ser Ser Cys Gly Ser Asn Thr Pro
269 415             420             425             430
271 tcg agt agt gat gat gtt gag gcg gat gca tca gaa agg caa gag gat     3484
272 Ser Ser Ser Asp Asp Val Glu Ala Asp Ala Ser Glu Arg Gln Glu Asp
273             435             440             445
275 ggc acc aat ggt gag gtg aaa gaa acg aat gaa gac act aat aaa cct     3532
276 Gly Thr Asn Gly Glu Val Lys Glu Thr Asn Glu Asp Thr Asn Lys Pro
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284 Thr Asp Pro Trp Lys Ser Val Ser Asp Glu
285     480                 485
287 tcaacttcct ttatttcaaa tcattttctc atataaatat tgtacattcg ggt cga     3686
288                                                         Gly Arg
289                                                             490
291 att gcc ttc caa gct ctc ttc tcc aga gag gta ttg ccg caa agt ttt   3734
292 Ile Ala Phe Gln Ala Leu Phe Ser Arg Glu Val Leu Pro Gln Ser Phe
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296 Thr Tyr Arg Glu Glu His Arg Glu Glu Glu Gln Gln Gln Gln Glu Gln
297             510                 515                 520
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300 Arg Tyr Pro Met Ala Leu Asp Leu Asn Phe Thr Ala Gln Leu Thr Pro
301         525                 530                 535
303 gtt gat gat caa gag gag aag aga aac aca gga ttt ctt gga atc gga   3878
304 Val Asp Asp Gln Glu Glu Lys Arg Asn Thr Gly Phe Leu Gly Ile Gly
305     540                 545                 550
307 tta gat gct tca aag cta atg agt aga gga aga aca ggt ttt aaa cca   3926
308 Leu Asp Ala Ser Lys Leu Met Ser Arg Gly Arg Thr Gly Phe Lys Pro
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311 tac aaa aga tgt tcc atg gaa gcc aaa gaa agt aga atc ctc aac aac   3974
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320 Glu Thr Gln Ala Ser Thr
321         605
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VERIFICATION SUMMARY   DATE: 07/31/2002
PATENT APPLICATION: US/10/084,553   TIME: 12:41:34

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